

Regioselectivity and Chemoselectivity Analysis of Oxazole and Thiazole Ring Formation by the Peptide-Heterocyclizing Microcin B17 Synthetase Using High-Resolution MS/MS

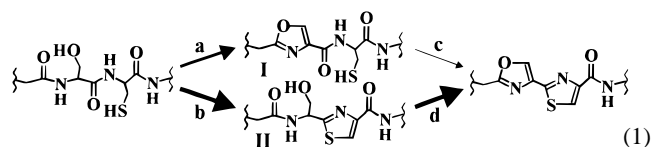
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The presence of rigid structural elements such as thiazole and oxazole heterocyclic rings in peptide-derived compounds confers a wide range of therapeutic properties including antibiotic, antiviral, and antitumor activity. Molecules with thiazoles and oxazoles in tandem 4,2-connected bisheterocyclic arrays are of particular interest, given the recent demonstration that the bithiazole moiety in the antitumor drug bleomycin is a DNA-targeting intercalator.¹ The tandem oxazole/thiazole and thiazole/oxazole pairs found in the *Escherichia coli* antibiotic microcin B17 (MccB17)² may target MccB17 to DNA and kill bacteria by the accumulation of double strand DNA breaks in a DNA gyrase dependent mechanism.³ Posttranslational ring biogenesis from Ser and Cys residues occurs on a complex of three proteins (McbB, C, D), utilizing Zn²⁺, FMN, and ATP to cyclize, dehydrate, and aromatize these substrate residues.⁴ Using high-resolution tandem mass spectrometry (MS/MS), we determine here the effect of heterocycle formation on MS/MS of reaction intermediates and use this unique MS/MS signature to ascertain the regio- and chemoselectivity of MccB17 synthetase.

The first heterocyclizable site in the 69 residue McbA substrate for posttranslational processing by MccB17 synthetase is the tripeptidyl unit Gly₃₉Ser₄₀Cys₄₁, converted to the tandem bisheterocycle.^{5a,b} Two rounds of cyclodehydration (−18.01 Da) and dehydrogenation (−2.02 Da) occur (eq 1) in both full length



substrate (McbA_{1–69}) and a McbA_{1–46} fragment. Discrete M-20 Da species accumulate in enzyme incubations, which then are converted to bisheterocycle.^{5b} These intermediates are presumed to be monocyclic species, and characterization would allow interrogation of preference of the bisheterocyclizing synthetase for regioselectivity and/or chemoselectivity in ring formation. McbA substrate recognition requirements^{5a,6} mandate that mechanistic investigations be conducted in at least the McbA_{1–46} substrate context (4.3 kDa). Substrates were generated as MBP–

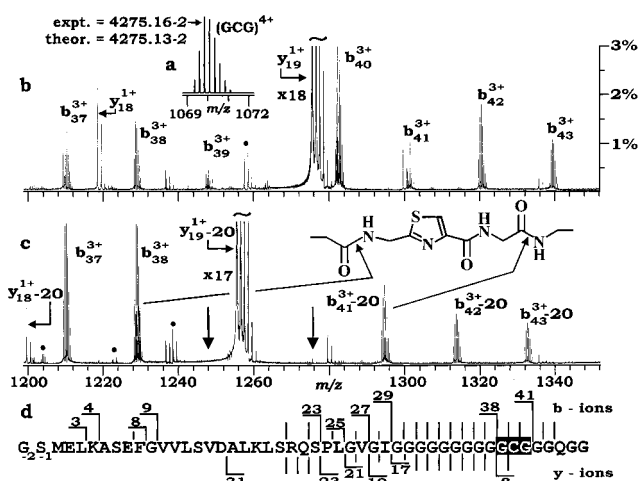


Figure 1. (a) Partial ESI/FT mass spectrum of the Gly₃₉Cys₄₀Gly₄₁ substrate (48 residues) of microcin B17 synthetase; 4+ ions, 10 scans, 9.4 T. (b) Partial MS/MS spectrum (*m/z* 1200–1350) from photodissociation of the Gly₃₉Cys₄₀Gly₄₁ substrate ions in (a); dot, H₂O loss peak; 500 scans, 9.4 T. (c) Partial MS/MS spectrum from photodissociation of the 4+ ions of the Gly thiazole–Gly enzymatic product; vertical arrows, theoretical positions of b₃₉³⁺ and (b₄₀–20)³⁺ fragment ions; dots, H₂O loss peaks; 500 scans, 9.4 T. (d) Correlation of all fragment ions observed in the full MS/MS spectrum of the (GCG)⁴⁺ ions with the primary sequence of this substrate; b- and y-type ions are indicated with vertical lines above and below the sequence, respectively.

McbA_{1–46} fusion proteins,⁵ subjected to MccB17 synthetase, and then cleaved at a thrombin site to release 48mer fragments GS–McbA_{1–46} (see Figure 1c) which were transferred into the gas phase by electrospray ionization (ESI)⁷ and analyzed in a 9.4^{8a} or 4.7^{8b} Tesla Fourier transform (FT) mass spectrometer.⁹

For MS/MS, it was anticipated that starting substrates could be sequenced partially via b-type (containing the N-terminus) and y-type (containing the C-terminus)¹⁰ fragment ions and that conversion of a peptide linkage into a thiazole or oxazole should abrogate cleavage at that site. As an initial test, a substrate mutant with only a single cyclizable residue, MBP–McbA_{1–46} G₃₉C₄₀G₄₁ (GCG), yielded multiply charged species with 3+, 4+, and 5+ charge states and the expected relative molecular weight (*M_r*) value (4275.16-2 vs 4275.13-2 Da,¹¹ theory; 7 ppm error). The (GCG)⁴⁺ ions (Figure 1a) were of highest abundance for subsequent isolation¹² and fragmentation,¹³ producing 20 y-type and 28 b-type fragment ions (Figure 1d) in the MS/MS spectrum. The b-ion series (b₃₇³⁺–b₄₃³⁺) between *m/z* 1200–1350 (Figure 1b) directly sequences the predicted site of thiazole formation. Conversion of G₃₉C₄₀G₄₁ to the amidomethylthiazole-G₄₁ product results in a loss of 20.02 Da ($\Delta m_{\text{theory}} = -20.03$ Da, data not shown); MS/MS of these (GCG–20)⁴⁺ ions reveals an absence of both the b₃₉³⁺ and the (b₄₀–20)³⁺ fragment ions (Figure 1c, vertical

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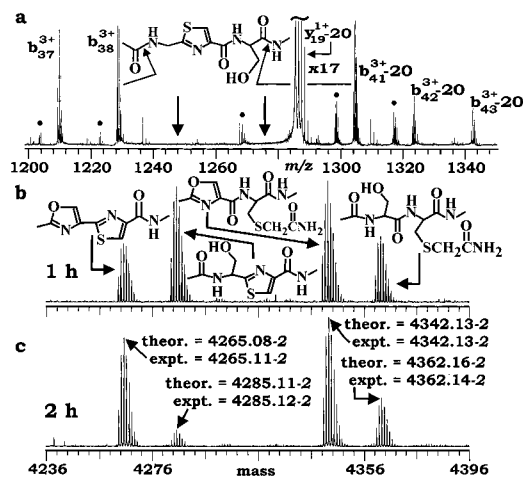


Figure 2. (a) Partial MS/MS spectrum (m/z 1200–1350) from photodissociation of $(M-20)^{4+}$ ions of the $\text{Gly}_{39}\text{Cys}_{40}\text{Ser}_{41}$ substrate; vertical arrows, theoretical positions of b_{39}^{3+} and $(b_{40}-20)^{3+}$ fragment ions; dots, H_2O loss peaks; 400 scans, 9.4 T. ESI/FT mass spectra (x -axis converted to mass) of the $\text{Gly}_{39}\text{Ser}_{40}\text{Cys}_{41}$ substrate after microcin synthetase processing for 1 h (b) and 2 h (c), with subsequent iodoacetamide treatment; 16 scans, 4.7 T.

arrows) while the b_{37} , b_{38} , $(b_{41}-20)^{3+}$, $(b_{42}-20)^{3+}$, and $(b_{43}-20)^{3+}$ ions are quite clear. Thus the monothiazole prevents fragmentation not only at the amide bond deleted by ring formation but also at the conjugated amide bond C-terminal to the ring. Another regioisomeric single site substrate, GGC, gave independent corroboration of this MS/MS effect, as in that case the b_{40}^{3+} and $(b_{41}-20)^{3+}$ ions were completely suppressed (>99% confidence) in the MS/MS spectrum of $(\text{GGC}-20)^{4+}$ ions (not shown). This heterocycle fingerprint now can be used analytically.

The regioisomeric double mutant $\text{Gly}_{39}\text{Cys}_{40}\text{Ser}_{41}$ (GCS), with Ser and Cys positionally interchanged from wild type McbA, has two potential sites for ring formation but has been observed to stall entirely at a single ring intermediate;^{5b} MS/MS of these $(\text{GCS}-20)^{4+}$ ions shows an absence of the b_{39}^{3+} and $(b_{40}-20)^{3+}$ fragment ions (Figure 2a), which are observed for the sample before enzymatic incubation (not shown). Thus the GCS-20 species is at least 97% thiazole; any oxazole-containing species would give rise to a b_{39} ion at ~ 1248 m/z (Figure 2a, vertical arrow). The thiolate anion (Cys S^-) is both more accessible than the alkoxide (Ser O^-) (pK_a 8 vs 13) and more nucleophilic, so that exclusive thiazole formation is the chemoselectivity. In addition, there may be a contributing regioselectivity programmed partially by the substrate local sequence, with cyclization to the amide carbonyl of glycine (no β -carbon) kinetically favored. The lack of any (<0.1%) bisheterocycle formation for this GCS substrate indicates that microcin synthetase in this sequence context cannot facilitate ring formation involving hydroxyl attack at an amide carbonyl that is in conjugation with a thiazole ring.

To equalize side chain reactivity and assess synthetase regioselectivity, the $\text{Gly}_{39}\text{Cys}_{40}\text{Cys}_{41}$ (GCC) and $\text{Gly}_{39}\text{Ser}_{40}\text{Ser}_{41}$ (GSS) mutant substrates were employed. The observed M_r values of the GCC- and the GSS-substrate were within 11 ppm of those expected, and MS/MS of the $4+$ ions of each gave the full b_{37}^{3+} – b_{43}^{3+} fragment ion series (not shown). The substrates were enzymatically processed to $\sim 95\%$ M-20 species in 8 h, with progression to M-40 bisheterocycles detectable ($\sim 5\%$) but very slow. In the MS/MS spectra of both the $(\text{GCC}-20)^{4+}$ and the $(\text{GSS}-20)^{4+}$ (Supporting Information), the b_{39}^{3+} and the $(b_{40}-20)^{3+}$ ions were not present, indicating that the intermediates are >90% and >92% N-terminal heterocycle, respectively. Even with equal intrinsic reactivities of two competing nucleophiles, the enzyme cyclizes mostly the N-terminal site, suggesting either that this first ring formation site is in register with a $\text{N} \rightarrow \text{C}$ directionality of processing and/or the presence of the upstream, sterically

permissive glycine has a significant effect on the cyclization rate. However, even with a thiolate as the intramolecular nucleophile, the enzyme does not catalyze efficient ring formation involving attack at an amide bond conjugated to a heterocyclic ring (<5% bithiazole in 8 h); therefore, a preferred $\text{N} \rightarrow \text{C}$ processing of bisheterocyclic sites appears unlikely. It may be that the C-terminal heterocycle (position 41) forms in the GCC- and GSS-substrates to a low level (<5%) and only this M-20 species is processed to the low level of observed M-40. This leads to a synthetase model wherein the M-20 intermediate with the N-terminal ring formed (path a, eq 1) is a kinetic “bottleneck” and this view is corroborated by the data below.

The wild type $\text{Gly}_{39}\text{Ser}_{40}\text{Cys}_{41}$ (GSC) sequence is the only substrate of this set to be processed to majority bisheterocyclic product. Its measured M_r value of 4305.16-2 Da is 5 ppm away from that expected (4305.14-2 Da), and MS/MS again yielded the diagnostic b_{37} – b_{43} ion series (not shown). MS/MS experiments of the $(\text{GSC}-20)^{4+}$ ions from reactions quenched at 1, 2, and 8 h showed that the b_{39} and b_{40} ions were absent in the 2 and 8 h data (indicating oxazole) but only the b_{40} fragment ion was completely suppressed in the 1 h data (not shown). This suggested that a mixture of oxazole (I) and thiazole (II) was present at early processing times. In principle, the oxazole/thiazole ratio could be determined by the $b_{41}-20/b_{39}$ ratio. However, iodoacetamide alkylation was used to separate I from II by adding 57.02 Da only to I with a remaining free cysteine thiol. For the partially processed, alkylated samples of GSC substrate at 1 h (Figure 2b), 2 h (Figure 2c), and 8 h (not shown), two conclusions can be drawn. First, both oxazole and thiazole monoheterocycles are formed (eq 1a and eq 1b followed) at a ratio of path a/b of about 0.6/1. Thus, the synthetase can equalize the effective nucleophilicity of Ser-OH and Cys-SH for cyclization on the upstream amide carbonyl. Second, the oxazole (I) apparently stalls at the M-20 stage while the thiazole (II) is processed on (path 1d) to M-40 bisheterocycle in ~ 2 h. By 8 h, a 25% progression of oxazole to bisheterocycle was measured, indicating a path c/d ratio of $\sim 0.06/1$. The enzyme has to balance chemoselectivity (nucleophilicity) and regioselectivity both in the first heterocyclization and second heterocyclization where β -substituents (Gly vs Cys/Ser vs heterocycle) and electrophilicity of the amide carbonyl differ. Thus, within this first bisheterocyclic site of McbA, processing to oxazole-thiazole occurs mainly with a $\text{C} \rightarrow \text{N}$ directionality.

Understanding the substrate requirements for efficient bisheterocycle production is critical in efforts to use the enzyme for combinatorial biosynthesis to generate combinations of tandem 4,2-heterocyclic systems for selection of new agents with biological activity. Future determination of regioselectivity of the synthetase toward substrates with up to eight sites of heterocycle formation will include MS/MS characterization of a second bisheterocyclization site, $\text{Gly}_{54}\text{Cys}_{55}\text{Ser}_{56}$ to see if indeed initial oxazole formation is the dominant kinetic pathway to the thiazole-oxazole pair.

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Supporting Information Available: Partial MS/MS spectrum from dissociation of $(\text{GCC}-20)^{4+}$ and $(\text{GSS}-20)^{4+}$ ions (2 pages, print/PDF). See any current masthead page for ordering information and Web access instructions.